

**AMENDMENTS TO THE CLAIMS**

1. (Original) A peptide selected from the following (a), (b), (c) or (d):

(a) a peptide consisting of the amino acid sequence as shown in SEQ ID NO: 4;

(b) a peptide which consists of the amino acid sequence as shown in SEQ ID NO: 4 having addition, deletion or substitution of one or a plurality of amino acids and has  $\beta$ -ionone ring-2-hydroxylase activity;

(c) a peptide which consists of an amino acid sequence having a 50% or more identity with the amino acid sequence as shown in SEQ ID NO: 4 and has  $\beta$ -ionone ring-2-hydroxylase activity;  
or

(d) a bacterium-derived peptide which is encoded by a DNA consisting of the nucleotide sequence as shown in SEQ ID NO: 3 or a DNA hybridizable to a complementary DNA to said DNA under stringent conditions and has  $\beta$ -ionone ring-2-hydroxylase activity.

2. (Original) A gene encoding a peptide selected from the following (a), (b), (c) or (d):

(a) a peptide consisting of the amino acid sequence as shown in SEQ ID NO: 4;

(b) a peptide which consists of the amino acid sequence as shown in SEQ ID NO: 4 having addition, deletion or substitution of one or a plurality of amino acids and has  $\beta$ -ionone ring-2-hydroxylase activity;

(c) a peptide which consists of an amino acid sequence having a 50% or more identity with the amino acid sequence as shown in SEQ ID NO: 4 and has  $\beta$ -ionone ring-2-hydroxylase activity;  
or

(d) a bacterium-derived peptide which is encoded by a DNA consisting of the nucleotide sequence as shown in SEQ ID NO: 3 or a DNA hybridizable to a complementary DNA to said DNA under stringent conditions and has  $\beta$ -ionone ring-2-hydroxylase activity.

3. (Original) A microorganism obtainable by introducing the gene according to claim 2 thereinto, wherein the microorganism is capable of introducing a hydroxyl group at the position 2 carbon of  $\beta$ -ionone ring.

4. (Original) A microorganism obtainable by introducing the gene according to claim 2 and other carotenoid biosynthesis genes thereinto, wherein the microorganism is capable of introducing a hydroxyl group at the position 2 carbon of  $\beta$ -ionone ring.

5. (Original) The microorganism according to claim 4, wherein the other carotenoid biosynthesis genes are all or a part of a gene cluster required for synthesizing  $\beta$ -ionone ring-containing carotenoids from farnesyl pyrophosphate.

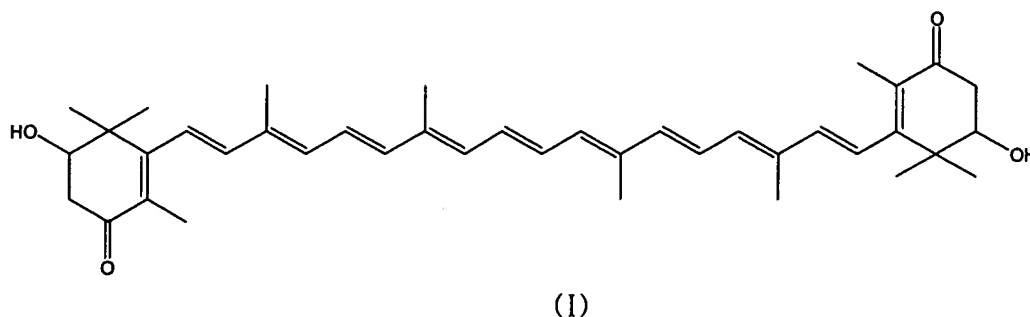
6. (Currently amended) The microorganism according to ~~any one of claims 3 to 5~~ claim 3, wherein the microorganism is *Escherichia coli*.

7. (Currently amended) A method of preparing a hydroxylated carotenoid, comprising culturing the microorganism according to ~~any one of claims 3 to 6~~ claim 3 in a medium and obtaining from

the resultant culture or cells a carotenoid which is hydroxylated at the position 2 carbon of its  $\beta$ -ionone ring.

8. (Original) The method according to claim 7, wherein the carotenoid which is hydroxylated at the position 2 carbon of its  $\beta$ -ionone ring is  $\beta,\beta$ -carotene-2-ol (2-hydroxy- $\beta$ -carotene),  $\beta,\beta$ -carotene-2,2'-diol (2,2'-dihydroxy- $\beta$ -carotene), caloxanthin (2-hydroxyzeaxanthin), nostoxanthin (2,2'-dihydroxyzeaxanthin), 2-hydroxy- $\beta,\beta$ -carotene-4,4'-dione (2-hydroxycanthaxanthin), 2,2'-dihydroxy- $\beta,\beta$ -carotene-4,4'-dione (2,2'-dihydroxycanthaxanthin), 2-hydroxyastaxanthin or 2,3,2',3'-tetrahydroxy- $\beta,\beta$ -carotene-4,4'-dione (2,2'-dihydroxyastaxanthin).

9. (Original) 2,2'-dihydroxy- $\beta,\beta$ -carotene-4,4'-dione (2,2'-dihydroxycanthaxanthin) represented by the following chemical formula (I):



10. (Original) An antioxidant comprising 2,2'-dihydroxy- $\beta,\beta$ -carotene-4,4'-dione (2,2'-dihydroxycanthaxanthin) or 2-hydroxy- $\beta,\beta$ -carotene-4,4'-dione (2-hydroxycanthaxanthin) as an active ingredient.

11. (Original) A gene encoding a peptide selected from the following (e), (f) or (g):
- (e) a peptide consisting of the amino acid sequence as shown in SEQ ID NO: 30;
  - (f) a peptide which consists of the amino acid sequence as shown in SEQ ID NO: 30 having addition, deletion or substitution of one or a plurality of amino acids and has  $\beta$ -ionone ring-3-hydroxylase activity; or
  - (g) a bacterium-derived peptide which is encoded by a DNA consisting of the nucleotide sequence as shown in SEQ ID NO: 29 or a DNA hybridizable to a complementary DNA to said DNA under stringent conditions and has  $\beta$ -ionone ring-3-hydroxylase activity.
12. (Original) A microorganism obtainable by introducing the gene according to claim 11 thereinto, wherein the microorganism is capable of introducing a hydroxyl group at the position 3 carbon of  $\beta$ -ionone ring.
13. (Original) A microorganism obtainable by introducing the gene according to claim 11 and other carotenoid biosynthesis genes thereinto, wherein the microorganism is capable of introducing a hydroxyl group at the position 3 carbon of  $\beta$ -ionone ring.
14. (Original) The microorganism according to claim 13, wherein the other carotenoid biosynthesis genes are all or a part of a gene cluster required for synthesizing  $\beta$ -ionone ring-containing carotenoids from farnesyl pyrophosphate.

15. (Currently amended) The microorganism according to ~~any one of claims 12 to 14~~ claim 12, wherein the microorganism is *Escherichia coli*.

16. (Currently amended) A method of preparing a hydroxylated carotenoid, comprising culturing the microorganism according to ~~any one of claims 12 to 15~~ claim 12 in a medium and obtaining from the resultant culture or cells a carotenoid which is hydroxylated at the position 3 carbon of its  $\beta$ -ionone ring.

17. (Original) A gene encoding a peptide selected from the following (h), (i) or (j):

(h) a peptide consisting of the amino acid sequence as shown in SEQ ID NO: 32;

(i) a peptide which consists of the amino acid sequence as shown in SEQ ID NO: 32 having addition, deletion or substitution of one or a plurality of amino acids and has geranylgeranyl pyrophosphate synthase activity; or

(j) a bacterium-derived peptide which is encoded by a DNA consisting of the nucleotide sequence as shown in SEQ ID NO: 31 or a DNA hybridizable to a complementary DNA to said DNA under stringent conditions and has geranylgeranyl pyrophosphate synthase activity.